

SEQUENCE LISTING

(1) GENERAL INFORMATION

- (i) APPLICANT: Nagai, Yosiyuki
Kato, Atsushi
Murai, Fukashi
Sakata, Tsuneaki
Sasegawa, Mamoru
Shioda, Tatsuo
- (ii) TITLE OF THE INVENTION: Recombinant Sendai Virus
- (iii) NUMBER OF SEQUENCES: 4
- (iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: Clark & Elbing LLP
(B) STREET: 176 Federal Street
(C) CITY: Boston
(D) STATE: MA
(E) COUNTRY: USA
(F) ZIP: 02110
- (v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Diskette
(B) COMPUTER: IBM Compatible
(C) OPERATING SYSTEM: DOS
(D) SOFTWARE: FastSEQ for Windows Version 2.0
- (vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER:
(B) FILING DATE:
(C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER: HEI 7-285417
(B) FILING DATE: 01-NOV-1995

(A) APPLICATION NUMBER: JP96/03069
(B) FILING DATE: 22-OCT-1996
- (viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Clark, Paul T
(B) REGISTRATION NUMBER: 30,162
(C) REFERENCE/DOCKET NUMBER: 50026/005001
- (ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: 617-428-0200
(B) TELEFAX: 617-428-7045
(C) TELEX:

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 38 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Other

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

TGCGGCCGCC GTACGGTGGC AATGAGTGAA GGAGAAGT

38

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 69 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Other

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TTGCGGCCGC GATGAAC TTT CACCCTAAGT TTTT VTTACT ACGGCGTACG TCATCTTTT
TCTCTCTGC

60

69

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Other

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

AAGCGGCCGC CAAAGTTCAC GATGGAAGAC

30

(2) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 69 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Other

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

TGCGGCCGCC ATGAACTTTC ACCCTAAGTT TTTCTTACTA CGGATTATTA CAATTGGAC
TTTCCGCCC

60
69